R. Li

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RAW SEQUENCE LISTING PATENT APPLICATION: US/09/727,739B

DATE: 06/06/2002 TIME: 15:01:04

Input Set : A:\020523.seq.list.txt

Output Set: N:\CRF3\06062002\I727739B.raw

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5 <110> APPLICANT: Sheridan, Mark
       Kittilson, Jeffrey
       Moore, Craig
13 <120> TITLE OF INVENTION: Somatostatins and Methods
17 <130> FILE REFERENCE: 255.00040101
21 <140> CURRENT APPLICATION NUMBER: US 09/727,739B
23 <141> CURRENT FILING DATE: 2000-12-01
27 <150> PRIOR APPLICATION NUMBER: US 60/168,934
29 <151> PRIOR FILING DATE: 1999-12-03
33 <160> NUMBER OF SEQ ID NOS: 52
37 <170> SOFTWARE: PatentIn version 3.0
41 <210> SEQ ID NO: 1
43 <211> LENGTH: 14
45 <212> TYPE: PRT
47 <213> ORGANISM: Homo sapiens, ...
51 <400> SEQUENCE: 1
53 Ala Gly Cys Lys Asn Phe Phe Trp Lys Thr Phe Thr Ser Cys
54 1 5
56 <210> SEQ ID NO: 2
58 <211> LENGTH: 14
60 <212> TYPE: PRT
62 <213> ORGANISM: Oncorhynchus mykiss
66 <400> SEQUENCE: 2
68 Ala Gly Cys Lys Asn Phe Tyr Trp Lys Gly Phe Thr Ser Cys
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.71 <210> SEQ ID NO: 3
73 <211> LENGTH: 114
75 <212> TYPE: PRT
77 <213> ORGANISM: Oncorhynchus mykiss
81 <400> SEQUENCE: 3
83 Met Leu Ser Thr Arg Val Gln Cys Ala Leu Ala Leu Leu Ser Leu Ala
                            10 15
84 1 . 5
86 Leu Ala Ile Ser Ser Val Ser Ala Ala Pro Ser Asp Ala Lys Leú Arg
                           25 30
87 20
89 Gln Leu Leu Gln Arg Ser Leu Met Ala Pro Ala Gly Lys Gln Glu Leu
90 35 40 45
92 Ala Arg Asn Thr Leu Val Glu Leu Leu Ser Glu Leu Ala His Val Glu
93 50 55
                             60
95 Asn Glu Ala Ile Glu Leu Asp Asp Met Ser His Gly Val Glu Gln Glu
                           75 80
                   70
98 Asp Val Asp Leu Glu Leu Glu Arg Ala Pro Gly Pro Val Leu Ala Pro
                                 90
101 Arg Glu Arg Lys Ala Gly Cys Lys Asn Phe Phe Trp Lys Thr Phe Thr
```

RAW SEQUENCE LISTING DATE: 06/06/2002 PATENT APPLICATION: US/09/727,739B TIME: 15:01:04

Input Set : A:\020523.seq.list.txt
Output Set: N:\CRF3\06062002\I727739B.raw

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105
                                                    110
               100
102
104 Ser Cys
107 <210> SEQ ID NO: 4
109 <211> LENGTH: 26
111 <212> TYPE: PRT
113 <213> ORGANISM: Oncorhynchus mykiss
117 <400> SEQUENCE: 4
119 Ala Pro Gly Pro Val Leu Ala Pro Arg Glu Arg Lys Ala Gly Cys Lys
      . 5
                                     10
122 Asn Phe Phe Trp Lys Thr Phe Thr Ser Cys
      20
                                 25
125 <210> SEQ ID NO: 5
127 <211> LENGTH: 88
129 <212> TYPE: PRT
131 <213> ORGANISM: Oncorhynchus mykiss
135 <400> SEQUENCE: 5
137 Met Leu Ser Thr Arg Val Gln Cys Ala Leu Ala Leu Leu Ser Leu Ala
                5
                                     10
140 Leu Ala Ile Ser Ser Val Ser Ala Ala Pro Ser Asp Ala Lys Leu Arg
     20
                                                  30
                                 25
141
143 Gln Leu Leu Gln Arg Ser Leu Met Ala Pro Ala Gly Lys Gln Glu Leu
144 35
                             40
146 Ala Arg Asn Thr Leu Val Glu Leu Leu Ser Glu Leu Ala His Val Glu
                                             60
147 50 ·
                          55
149 Asn Glu Ala Ile Glu Leu Asp Asp Met Ser His Gly Val Glu Gln Glu
                     70
150 65
152 Asp Val Asp Leu Glu Leu Glu Arg
153
                   85
155 <210> SEQ ID NO: 6
157 <211> LENGTH: 12
159 <212> TYPE: PRT
161 <213> ORGANISM: Oncorhynchus mykiss
165 <400> SEQUENCE: 6
167 Ala Pro Gly Pro Val Leu Ala Pro Arg Glu Arg Lys
168 1
                   5
                       10
170 <210> SEQ ID NO: 7
172 <211> LENGTH: 24
174 <212> TYPE: PRT
176 <213> ORGANISM: Oncorhynchus mykiss
180 <400> SEQUENCE: 7
182 Met Leu Ser Thr Arg Val Gln Cys Ala Leu Ala Leu Leu Ser Leu Ala
183 1
                                      10
185 Leu Ala Ile Ser Ser Val Ser Ala
186
               20
188 <210> SEQ ID NO: 8
190 <211> LENGTH: 763
192 <212> TYPE: DNA
194 <213> ORGANISM: Oncorhynchus mykiss
198 <400> SEQUENCE: 8
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/727,739B

DATE: 06/06/2002 TIME: 15:01:04

Input Set : A:\020523.seq.list.txt

Output Set: N:\CRF3\06062002\I727739B.raw

```
199 ggggggggg gaacaggagc agcagaactc aaagagaagc caatctcaac gattgtctgc
201 ccaattgaac caccittatc catcctctgc ctcccccgag acccagaaga agatgctctc
                                                                       120
                                                                       180
203 gacgcgtgtc cagtgcgccc tagcactact ctccctagcc ctggccatca gcagcgtctc
                                                                       240
205 tgccgctccg tccgatgcca aactccgcca gctgctccaa cggtcactca tggcacctgc
                                                                       300
207 aggcaaacag gagcttgcca ggaatacact cgtagagcta ctctcagagc tcgcacatgt
209 agagaacgag gcgattgaat tggatgacat gtctcatggc gtggagcagg aggatgtgga
                                                                       360
211 tetegagetg gagegtgeae eeggeecagt actggeteea egtgaaegea aggetggatg
                                                                       420
213 caagaacttc ttctggaaga cctttacatc gtgttaatga atctactcct ttactgtgtg
                                                                       480
215 tactacatct catctcttt gtttcaatca ctcattgctg aatccaatgc accatggcct
                                                                       540
217 aaccetecte tteaaaaaat ttaaataaae aetgttataa etttaaeaat eattetgatg
                                                                       600
                                                                        660
219 tttctatcgc tcacttagat ttttttccga aaaggaacac aagaaagaat gttctacaaa
221 tgtatgcggt tctgctttga ctgtgattta tgtattttgg cagactattt ttaattgttt
                                                                       720
                                                                        763
226 <210> SEQ ID NO: 9
228 <211> LENGTH: 115
230 <212> TYPE: PRT
232 <213> ORGANISM: Oncorhynchus mykiss
236 <400> SEQUENCE: 9
238 Met Lys Val Cys Arg Ile His Cys Ala Leu Ala Leu Leu Gly Leu Ala
                 5
                                       10
239 1
241 Leu Ala Ile Cys Ser Gln Gly Ala Ala Ser Gln Pro Asp Leu Asp Leu
                                   25
      • .
               20
242
244 Arg Ser Arg Arg Leu Leu Gln Arg Ala Arg Ala Ala Leu Pro His
            35
                               40
245
247 Arg Ser Gly Val Ser Glu Arg Trp Arg Thr Phe Tyr Pro Asn Cys Pro
                           55
     50
250 Cys Leu Arg Pro Arg Lys Val Lys Cys Pro Ala Gly Ala Lys Glu Asp
                       70
                                           75
251 65
253 Leu Arg Val Glu Leu Glu Arg Ser Val Gly Asn Pro Asn Asn Leu Pro
                                       90
256 Pro Arg Glu Arg Lys Ala Gly Cys Lys Asn Phe Tyr Trp Lys Gly Phe
                                                       110
257
                100
                                   105
259 Thr Ser Cys
260
            115
262 <210> SEQ ID NO: 10
264 <211> LENGTH: 28
266 <212> TYPE: PRT
268 <213> ORGANISM: Oncorhynchus mykiss
272 <400> SEQUENCE: 10
274 Ser Val Gly Asn Pro Asn Asn Leu Pro Pro Arg Glu Arg Lys Ala Gly
                   5
                                       10
275 1
277 Cys Lys Asn Phe Tyr Trp Lys Gly Phe Thr Ser Cys
278
               20
280 <210> SEQ ID NO: 11
282 <211> LENGTH: 87
284 <212> TYPE: PRT
286 <213> ORGANISM: Oncorhynchus mykiss
290 <400> SEQUENCE: 11
292 Met Lys Val Cys Arg Ile His Cys Ala Leu Ala Leu Leu Gly Leu Ala
```

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/727,739B

DATE: 06/06/2002 TIME: 15:01:04

Input Set : A:\020523.seq.list.txt

Output Set: N:\CRF3\06062002\I727739B.raw

```
10
                    5
295 Leu Ala Ile Cys Ser Gln Gly Ala Ala Ser Gln Pro Asp Leu Asp Leu
                                25
                                                        30
                20
298 Arg Ser Arg Arg Leu Leu Gln Arg Ala Arg Ala Ala Leu Pro His
           35
                               40
                                                   45
301 Arg Ser Gly Val Ser Glu Arg Trp Arg Thr Phe Tyr Pro Asn Cys Pro
                           55
                                              ~ 60
       50
304 Cys Leu Arg Pro Arg Lys Val Lys Cys Pro Ala Gly Ala Lys Glu Asp
                       70
305 65
307 Leu Arg Val Glu Leu Glu Arg
310 <210> SEQ ID NO: 12
312 <211> LENGTH: 14
314 <212> TYPE: PRT
316 <213> ORGANISM: Oncorhynchus mykiss
320 <400> SEQUENCE: 12
322 Ser Val Gly Asn Pro Asn Asn Leu Pro Pro Arg Glu Arg Lys
                                        10
325 <210> SEQ ID NO: 13
327 <211> LENGTH: 25
329 <212> TYPE: PRT
331 <213> ORGANISM: Oncorhynchus mykiss
335 <400> SEQUENCE: 13
337 Met Lys Val Cys Arg Ile His Cys Ala Leu Ala Leu Leu Gly Leu Ala
                5
338 1
340 Leu Ala Ile Cys Ser Gln Gly Ala Ala
                20
341
343 <210> SEQ ID NO: 14
345 <211> LENGTH: 623
347 <212> TYPE: DNA
349 <213> ORGANISM: Oncorhynchus mykiss
353 <400> SEQUENCE: 14
354 accaggectg etecataceg actgatecag ategageata geceggteea geteageteg
356 teteacegeg tgccatecet gcaaacaaaa eccagetetg ttggagatga aggtetgeeg
                                                                         180
358 aatccactgt gccctggccc tgctgggttt ggccctggcc atttgcagcc aaggagccgc
360 ctcgcagccc gacctggacc tccgcagccg cagactcctt cagagggctc gtgccgctgc
                                                                         240
                                                                         300
362 attgccacac aggagtggag taagcgagcg gtggaggaca ttctatccca actgtccttg
364 cctgaggccc aggaaagtga agtgtcaagc gggggctaaa gaggacctgc gtgtggagct
                                                                         360
366 ggagcgctca gtgggcaacc ccaacaacct tcccccccgt gagcgcaaag ccggctgcaa
                                                                         420
368 gaacttctac tggaagggct tcacttcctg ctgagggaag aataaaccga ccaccttatg
                                                                         480
                                                                         540
370 acatgacgct gccaatcacg tcacaccgcc aacttacacc tgacgaatgc agccaatcaa
372 cagttagctg tgcccgatga tggttcttga aatcaacaga atgatgtacc tgtctaattt
                                                                         600
                                                                         623
374 gtgaaataaa tataaaataa ttg
377 <210> SEQ ID NO: 15
379 <211> LENGTH: 111
381 <212> TYPE: PRT
383 <213> ORGANISM: Oncorhynchus mykiss
387 <400> SEQUENCE: 15
389 Met Arg Val Ser Gln Ile His Cys Ala Leu Ala Leu Leu Gly Leu Ala
```

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/727,739B

DATE: 06/06/2002 TIME: 15:01:04

Input Set : A:\020523.seq.list.txt
Output Set: N:\CRF3\06062002\I727739B.raw

```
10
390 1
392 Leu Ala Ile Cys Ser Gln Gly Ala Ala Ser Gln Pro Asp Leu Asp Leu
              20
                                25
395 Ala Ser Arg Arg Leu Leu Gln Arg Ala Leu Ala Ala Leu Pro His
                             40
                                                45
398 Arg Ser Gly Val Ser Glu Arg Trp Arg Thr Phe Tyr Pro Asn Cys Pro
                                            60
401 Cys Leu Arg Trp Arg Pro Arg Lys Val Lys Gly Pro Gln Leu Lys Ala
                                        75
                      70
                         1 .
402 65
404 Lys Glu Asp Leu Glu Arg Ser Val Asp Asn Leu Pro Pro Arg Glu Arg
                 85
                                    90
407 Lys Ala Gly Cys Lys Asn Phe Tyr Trp Lys Gly Phe Thr Ser Cys
                        105
      100
408
410 <210> SEQ ID NO: 16
412 <211> LENGTH: 25
414 <212> TYPE: PRT
416 <213> ORGANISM: Oncorhynchus mykiss
420 <400> SEQUENCE: 16
422 Ser Val Asp Asn Leu Pro Pro Arg Glu Arg Lys Ala Gly Cys Lys Asn
423 1 5 10
425 Phe Tyr Trp Lys Gly Phe Thr Ser Cys
              20
428 <210> SEQ ID NO: 17
430 <211> LENGTH: 86
432 <212> TYPE: PRT
434 <213> ORGANISM: Oncorhynchus mykiss
438 <400> SEQUENCE: 17
440 Met Arg Val Ser Gln Ile His Cys Ala Leu Ala Leu Leu Gly Leu Ala
              5
                                     10
441 1
443 Leu Ala Ile Cys Ser Gln Gly Ala Ala Ser Gln Pro Asp Leu Asp Leu
444
              20
                                 25
446 Ala Ser Arg Arg Leu Leu Gln Arg Ala Leu Ala Ala Leu Pro His
447 35
                  40
449 Arg Ser Gly Val Ser Glu Arg Trp Arg Thr Phe Tyr Pro Asn Cys Pro
                          55
                                             60
       50
452 Cys Leu Arg Trp Arg Pro Arg Lys Val Lys Gly Pro Gln Leu Lys Ala
                      70
                                         75
453 65
455 Lys Glu Asp Leu Glu Arg
4.56
458 <210> SEQ ID NO: 18
460 <211> LENGTH: 11
462 <212> TYPE: PRT
464 <213> ORGANISM: Oncorhynchus mykiss
468 <400> SEQUENCE: 18
470 Ser Val Asp Asn Leu Pro Pro Arg Glu Arg Lys
471 1
473 <210> SEQ ID NO: 19
475 <211> LENGTH: 25
477 <212> TYPE: PRT
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RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/09/727,739B

DATE: 06/06/2002 TIME: 15:01:05

Input Set : A:\020523.seq.list.txt

Output Set: N:\CRF3\06062002\I727739B.raw

## Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:24; N Pos. 36,37,41,42,46,47

## Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seg#:22,23,24,25,26,33,34,50,51,52

## VERIFICATION SUMMARY

PATENT APPLICATION: US/09/727,739B

DATE: 06/06/2002 TIME: 15:01:05

Input Set : A:\020523.seq.list.txt

Output Set: N:\CRF3\06062002\I727739B.raw

L:547 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:22 L:565 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:23 L:583 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:24 L:622 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0 L:631 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:25 L:649 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:26 L:772 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:33 L:790 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:34 L:805 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (35) SEQUENCE: L:1308 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:50 L:1326 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:51 L:1344 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:52